

1/55

<110> Ben-Sasson, Shmuel A.

<120> Short Peptides Which Selectively Modulate the Activity of Protein Kinases

<130> 1242.1029-000 (CMCC-679)

<140> US 09/161,094

<141> 1998-09-25

<160> 172

<170> FastSEQ for Windows Version 4.0

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<213> unknown

<220>

<223> c-Raf

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Thr Gln Trp Cys Glu Gly Ser Ser Leu Tyr Lys His Leu His Val Gln
1 5 10 15
Glu Thr Lys Phe
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<213> unknown

$\langle 220 \rangle$

<223> a-Raf

<400> 2

Thr Gln Trp Cys Glu Gly Ser Ser Leu Tyr His His Leu His Val Ala
1 5 10 15
Asp Thr Arg Phe
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<220>

<223> Braf

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Thr Gln Trp Cys Glu Gly Ser Ser Leu Tyr His His Leu His Ile Ile
1 5 10 15
Glu Thr Lys Phe

20

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 1 5 10 15
 Gly Arg Phe

<210> 5
 <211> 19
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 <213> unknown

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 Met Glu Tyr Val Pro Gly Gly Glu Met Phe Ser Arg Leu Gln Arg Val
 1 5 10 15
 Gly Arg Phe

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<400> 6
 Met Glu Tyr Val Asn Gly Gly Asp Leu Met Tyr His Ile Gln Gln Val
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 Gly Lys Phe

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 <213> unknown

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 Met Glu Tyr Val Asn Gly Gly Asp Leu Met Tyr His Ile Gln Gln Val
 1 5 10 15

Gly Arg Phe

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 <211> 19
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 <213> unknown

<220>
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 Met Glu Tyr Val Thr Gly Gly Asp Leu Met Tyr His Ile Gln Gln Leu
 1 5 10 15
 Gly Lys Phe

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 1 5 10 15
 Gly Arg Phe

<210> 10
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 <212> PRT
 <213> unknown

<220>
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 Met Glu Tyr Val Asn Gly Gly Asp Leu Met Phe Gln Ile Gln Arg Ser
 1 5 10 15
 Arg Lys Phe

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<220>
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Arg Arg Phe

5

10

15

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Met Glu Tyr Leu Asn Gly Gly Asp Leu Met Tyr His Ile Gln Ser Cys
1 5 10 15
His Lys Phe

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<211> 19
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<213> unknown

<220>
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<400> 13

Met Glu Tyr Ala Asn Gly Gly Glu Leu Phe Phe His Leu Ser Arg Glu
1 5 10 15
Arg Val Phe

<210> 14
<211> 22
<212> PRT
<213> unknown

<220>
<223> GSK3a

<400> 14

Leu Glu Tyr Val Pro Glu Thr Val Tyr Arg Val Ala Arg His Phe Thr
1 5 10 15
Lys Ala Lys Leu Ile Ile
20

<210> 15
<211> 22
<212> PRT
<213> unknown

<220>
<223> GSK3b

<400> 15

Leu	Asp	Tyr	Val	Pro	Glu	Thr	Val	Tyr	Arg	Val	Ala	Arg	His	Tyr	Ser
1				5					10					15	
Arg	Ala	Lys	Gln	Thr	Leu										
			20												

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<211> 16

<212> PRT

<213> unknown

<220>

<223> CK IIa

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Phe	Glu	His	Val	Asn	Asn	Thr	Asp	Phe	Lys	Gln	Leu	Tyr	Gln	Thr	Leu
1				5					10					15	

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<211> 16

<212> PRT

<213> unknown

<220>

<223> CK IIa'

<400> 17

Phe	Glu	Tyr	Ile	Asn	Asn	Thr	Asp	Phe	Lys	Gln	Leu	Tyr	Gln	Ile	Leu
1				5					10					15	

<210> 18

<211> 19

<212> PRT

<213> unknown

<220>

<223> bARK1

<400> 18

Leu	Asp	Leu	Met	Asn	Gly	Gly	Asp	Leu	His	Tyr	His	Leu	Ser	Gln	His
1				5					10					15	
Gly	Val	Phe													

<210> 19

<211> 23

<212> PRT

<213> unknown

<220>

<223> GRK1

<400> 19

Met	Thr	Ile	Met	Asn	Gly	Gly	Asp	Ile	Arg	Tyr	His	Ile	Tyr	Asn	Val
-----	-----	-----	-----	-----	-----	-----	-----	-----	-----	-----	-----	-----	-----	-----	-----

1	5	10	15
Asp	Glu	Asp	Asn
	Pro	Gly	Phe
	20		

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<220>
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1				5					10					15
Gly	Asn	Pro	Gly	Phe										
			20											

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 <213> unknown

<220>
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Leu	Thr	Ile	Met	Asn	Gly	Gly	Asp	Leu	Lys	Phe	His	Ile	Tyr	Asn
1				5					10					15
Gly	Asn	Pro	Gly	Phe										
			20											

<210> 22
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 <212> PRT
 <213> unknown

<220>
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<400> 22														
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1				5					10					15
Gly	Gln	Ala	Gly	Phe										
			20											

<210> 23
 <211> 19
 <212> PRT
 <213> unknown

<220>
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<400> 23

Met Gln Leu Val Ser Gly Gly Glu Leu Phe Asp Arg Ile Val Glu Lys
 1 5 10 15
 Gly Gly Tyr

<210> 24
 <211> 19
 <212> PRT
 <213> unknown

<220>
 <223> CaMK IIa

<400> 24

Phe Asp Leu Val Thr Gly Gly Glu Leu Phe Glu Asp Ile Val Ala Arg
 1 5 10 15
 Glu Tyr Tyr

<210> 25
 <211> 19
 <212> PRT
 <213> unknown

<220>
 <223> Plk

<400> 25

Leu Glu Leu Cys Arg Arg Arg Ser Leu Leu Glu Leu His Lys Arg Arg
 1 5 10 15
 Lys Ala Leu

<210> 26
 <211> 19
 <212> PRT
 <213> unknown

<220>
 <223> Plx1

<400> 26

Leu Glu Leu Cys Arg Arg Arg Ser Leu Leu Glu Leu His Lys Arg Arg
 1 5 10 15
 Lys Ala Val

<210> 27
 <211> 19
 <212> PRT
 <213> unknown

<220>
 <223> POLO

8/55

<400> 27

Leu Glu Leu Cys Lys Lys Arg Ser Met Met Glu Leu His Lys Arg Arg
1 5 10 15
Lys Ser Ile

<210> 28

<211> 19

<212> PRT

<213> unknown

<220>

<223> SNK

<400> 28

Leu Glu Tyr Cys Ser Arg Arg Ser Met Ala His Ile Leu Lys Ala Arg
1 5 10 15
Lys Val Leu

<210> 29

<211> 19

<212> PRT

<213> unknown

<220>

<223> CDC 5

<400> 29

Leu Glu Ile Cys Pro Asn Gly Ser Leu Met Glu Leu Leu Lys Arg Arg
1 5 10 15
Lys Val Leu

<210> 30

<211> 20

<212> PRT

<213> unknown

<220>

<223> Sak

<400> 30

Leu Glu Met Cys His Asn Gly Glu Met Asn Arg Tyr Leu Lys Asn Arg
1 5 10 15
Val Lys Pro Phe
20

<210> 31

<211> 19

<212> PRT

<213> unknown

<220>

<223> Prk

9/55

<400> 31

Leu Glu Leu Cys Ser Arg Lys Ser Leu Ala His Ile Trp Lys Ala Arg
1 5 10 15
His Thr Leu

<210> 32

<211> 19

<212> PRT

<213> unknown

<220>

<223> Plol

<400> 32

Leu Glu Leu Cys Glu His Lys Ser Leu Met Glu Leu Leu Arg Lys Arg
1 5 10 15
Lys Gln Leu

<210> 33

<211> 19

<212> PRT

<213> unknown

<220>

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<400> 33

Met Glu Tyr Ala Ser Gly Gly Glu Val Phe Asp Tyr Leu Val Ala His
1 5 10 15
Gly Arg Met

<210> 34

<211> 19

<212> PRT

<213> unknown

<220>

<223> P78

<400> 34

Met Glu Tyr Ala Ser Gly Gly Glu Val Phe Asp Tyr Leu Val Ala His
1 5 10 15
Gly Arg Met

<210> 35

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<212> PRT

<213> unknown

<220>

<223> CDK2

<400> 35

Phe	Glu	Phe	Leu	His	Gln	Asp	Leu	Lys	Lys	Phe	Met	Asp	Ala	Ser	Ala
1				5					10					15	
Leu	Thr	Gly	Ile												
			20												

<210> 36

<211> 20

<212> PRT

<213> unknown

<220>

<223> CDK4

<400> 36

Phe	Glu	His	Val	Asp	Gln	Asp	Leu	Arg	Thr	Tyr	Leu	Asp	Lys	Ala	Pro
1				5					10					15	
Pro	Pro	Gly	Leu												
			20												

<210> 37

<211> 20

<212> PRT

<213> Unknown

<220>

<223> CDK6

<400> 37

Phe	Glu	His	Val	Asp	Gln	Asp	Leu	Thr	Thr	Tyr	Leu	Asp	Lys	Val	Pro
1				5					10					15	
Glu	Pro	Gly	Val												
			20												

<210> 38

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<212> PRT

<213> unknown

<220>

<223> c-Src

<400> 38

Thr	Glu	Tyr	Met	Ser	Lys	Gly	Ser	Leu	Leu	Asp	Phe	Leu	Lys	Gly	Glu
1				5					10					15	
Thr	Gly	Lys	Tyr	Leu											
			20												

<210> 39

<211> 21

<212> PRT

<213> unknown

<220>

<223> c-Yes

<400> 39

Thr	Glu	Phe	Met	Ser	Lys	Gly	Ser	Leu	Leu	Asp	Phe	Leu	Lys	Glu	Gly
1				5					10					15	
Asp	Gly	Lys	Tyr	Leu											
				20											

<210> 40

<211> 21

<212> PRT

<213> unknown

<220>

<223> Fyn

<400> 40

Thr	Glu	Tyr	Met	Asn	Lys	Gly	Ser	Leu	Leu	Asp	Phe	Leu	Lys	Asp	Gly
1				5					10					15	
Glu	Gly	Arg	Ala	Leu											
				20											

<210> 41

<211> 21

<212> PRT

<213> unknown

<220>

<223> c-Fgr

<400> 41

Thr	Glu	Phe	Met	Cys	His	Gly	Ser	Leu	Leu	Asp	Phe	Leu	Lys	Asn	Pro
1				5					10					15	
Glu	Gly	Gln	Asp	Leu											
				20											

<210> 42

<211> 21

<212> PRT

<213> unknown

<220>

<223> Lyn

<400> 42

Thr	Glu	Tyr	Met	Ala	Lys	Gly	Ser	Leu	Leu	Asp	Phe	Leu	Lys	Ser	Asp
1				5					10					15	
Glu	Gly	Gly	Lys	Val											
				20											

<210> 43

<211> 21

<212> PRT

<213> unknown

<220>

<223> Hck

<400> 43

Thr	Glu	Phe	Met	Ala	Lys	Gly	Ser	Leu	Leu	Asp	Phe	Leu	Lys	Ser	Asp
1				5				10						15	
Glu	Gly	Ser	Lys	Gln											
			20												

<210> 44

<211> 21

<212> PRT

<213> unknown

<220>

<223> Lck

<400> 44

Thr	Glu	Tyr	Met	Glu	Asn	Gly	Ser	Leu	Val	Asp	Phe	Leu	Lys	Thr	Pro
1				5				10						15	
Ser	Gly	Ile	Lys	Leu											
			20												

<210> 45

<211> 21

<212> PRT

<213> unknown

<220>

<223> Csk

<400> 45

Thr	Glu	Tyr	Met	Ala	Lys	Gly	Ser	Leu	Val	Asp	Tyr	Leu	Arg	Ser	Arg
1				5				10						15	
Gly	Arg	Ser	Val	Leu											
			20												

<210> 46

<211> 21

<212> PRT

<213> unknown

<220>

<223> MatK

<400> 46

Met	Glu	His	Val	Ser	Lys	Gly	Asn	Leu	Val	Asn	Phe	Leu	Arg	Thr	Arg
1				5				10						15	
Gly	Arg	Ala	Leu	Val											
			20												

<210> 47

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 <212> PRT
 <213> unknown

<220>
 <223> Fak

<400> 47
 Met Glu Leu Cys Thr Leu Gly Glu Leu Arg Ser Phe Leu Gln Val Arg
 1 5 10 15
 Lys Tyr Ser Leu
 20

<210> 48
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 <212> PRT
 <213> unknown

<220>
 <223> c-Abl

<400> 48
 Thr Glu Phe Met Thr Tyr Gly Asn Leu Leu Asp Tyr Leu Arg Glu Cys
 1 5 10 15
 Asn Arg Gln Glu Val
 20

<210> 49
 <211> 35
 <212> PRT
 <213> unknown

<220>
 <223> Tie

<400> 49
 Ile Glu Tyr Ala Pro Tyr Gly Asn Leu Leu Asp Phe Leu Arg Lys Ser
 1 5 10 15
 Arg Val Leu Glu Thr Asp Pro Ala Phe Ala Arg Glu His Gly Thr Ala
 20 25 30
 Ser Thr Leu
 35

<210> 50
 <211> 35
 <212> PRT
 <213> unknown

<220>
 <223> Tek

<400> 50
 Ile Glu Tyr Ala Pro His Gly Asn Leu Leu Asp Phe Leu Arg Lys Ser
 1 5 10 15

14/55

Arg Val Leu Glu Thr Asp Pro Ala Phe Ala Ile Ala Asn Ser Thr Ala
20 25 30
Ser Thr Leu
35

<210> 51
<211> 35
<212> PRT
<213> unknown

<220>
<223> Flg

<400> 51
Val Glu Tyr Ala Ser Lys Gly Asn Leu Arg Glu Tyr Leu Gln Ala Arg
1 5 10 15
Arg Pro Pro Gly Leu Glu Tyr Cys Tyr Asn Pro Ser His Asn Pro Glu
20 25 30
Glu Gln Leu
35

<210> 52
<211> 35
<212> PRT
<213> unknown

<220>
<223> Bek

<400> 52
Val Glu Tyr Ala Ser Lys Gly Asn Leu Arg Glu Tyr Leu Arg Ala Arg
1 5 10 15
Arg Pro Pro Gly Met Glu Tyr Ser Tyr Asp Ile Asn Arg Val Pro Glu
20 25 30
Glu Gln Met
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<210> 53
<211> 35
<212> PRT
<213> unknown

<220>
<223> FGFR-3

<400> 53
Val Glu Tyr Ala Ala Lys Gly Asn Leu Arg Glu Phe Leu Arg Ala Arg
1 5 10 15
Arg Pro Pro Gly Leu Asp Tyr Ser Phe Asp Thr Cys Lys Pro Pro Glu
20 25 30
Glu Gln Leu
35

<210> 54

<211> 35
 <212> PRT
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<400> 54
 Val Glu Cys Ala Ala Lys Gly Asn Leu Arg Glu Phe Leu Arg Ala Arg
 1 5 10 15
 Arg Pro Pro Gly Pro Asp Leu Ser Pro Asp Gly Pro Arg Ser Ser Glu
 20 25 30
 Gly Pro Leu
 35

<210> 55
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 <212> PRT
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<400> 55
 Thr Glu Tyr Cys Phe Tyr Gly Asp Leu Val Asn Tyr Leu His Lys Asn
 1 5 10 15
 Arg Asp Ser Phe Leu Ser His His Pro Glu Lys Pro Lys Lys Glu Leu
 20 25 30
 Asp Ile Phe Gly Leu Asn Pro Ala
 35 40

<210> 56
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 <212> PRT
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<220>
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<400> 56
 Thr Glu Tyr Cys Arg Tyr Gly Asp Leu Val Asp Tyr Leu His Arg Asn
 1 5 10 15
 Lys His Thr Phe Leu Gln His His Ser Asp Lys Arg Arg Pro Pro Ser
 20 25 30
 Ala Glu Leu Tyr Ser Asn Ala Leu
 35 40

<210> 57
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 <212> PRT
 <213> unknown

<220>

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Leu Pro Tyr Met Lys His Gly Asp Leu Arg Asn Phe Ile Arg Asn Glu
1 5 10 15
Thr His Asn Pro
20

<210> 61
 <211> 20
 <212> PRT
 <213> unknown

<220>
 <223> c-Sea

<400> 61
 Leu Pro Tyr Met Arg His Gly Asp Leu Arg His Phe Ile Arg Ala Gln
 1 5 10 15
 Glu Arg Ser Pro
 20

<210> 62
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 <212> PRT
 <213> unknown

<220>
 <223> Ron

<400> 62
 Leu Pro Tyr Met Cys His Gly Asp Leu Leu Gln Phe Ile Arg Ser Pro
 1 5 10 15
 Gln Arg Asn Pro
 20

<210> 63
 <211> 20
 <212> PRT
 <213> unknown

<220>
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<400> 63
 Thr Gln Leu Met Pro Phe Gly Cys Leu Leu Asp Tyr Val Arg Glu His
 1 5 10 15
 Lys Asp Asn Ile
 20

<210> 64
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<220>
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<400> 64
 Thr Gln Leu Met Pro Tyr Gly Cys Leu Leu Asp His Val Arg Glu Asn
 1 5 10 15
 Arg Gly Arg Leu

20

<210> 65
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 <212> PRT
 <213> unknown

<220>
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<400> 65
 Thr Gln Tyr Leu Pro Leu Gly Ser Leu Leu Asp His Val Arg Gln His
 1 5 10 15
 Arg Gly Ala Leu
 20

<210> 66
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 <212> PRT
 <213> unknown

<220>
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<400> 66
 Thr Gln Leu Met Pro His Gly Cys Leu Leu Glu Tyr Val His Glu His
 1 5 10 15
 Lys Asp Asn Ile
 20

<210> 67
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 <213> unknown

<220>
 <223> Ret

<400> 67
 Val Glu Tyr Ala Lys Tyr Gly Ser Leu Arg Gly Phe Leu Arg Glu Ser
 1 5 10 15
 Arg Lys Val Gly Pro Gly Tyr Leu Gly Ser Gly Gly Ser Arg Asn Ser
 20 25 30
 Ser Ser Leu Asp His Pro Asp Glu Arg Ala Leu
 35 40

<210> 68
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19/55

<400> 68

Phe Glu Tyr Met Arg His Gly Asp Leu Asn Arg Phe Leu Arg Ser His
1 5 10 15
Gly Pro Asp Ala Lys Leu Leu Ala Gly Gly Glu Asp Val Ala Pro Gly
20 25 30
Pro Leu

<210> 69

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<212> PRT

<213> unknown

<220>

<223> TrkB

<400> 69

Phe Glu Tyr Met Lys His Gly Asp Leu Asn Lys Phe Leu Arg Ala His
1 5 10 15
Gly Pro Asp Ala Val Leu Met Ala Glu Gly Asn Pro Pro Thr Glu Leu
20 25 30

<210> 70

<211> 35

<212> PRT

<213> unknown

<220>

<223> TrkC

<400> 70

Phe Glu Tyr Met Lys His Gly Asp Leu Asn Lys Phe Leu Arg Ala His
1 5 10 15
Gly Pro Asp Ala Met Ile Leu Val Asp Gly Gln Pro Arg Gln Ala Lys
20 25 30
Gly Glu Leu
35

<210> 71

<211> 19

<212> PRT

<213> unknown

<220>

<223> Syk

<400> 71

Met Glu Met Ala Glu Leu Gly Pro Leu Asn Lys Tyr Leu Gln Gln Asn
1 5 10 15
Arg His Val

<210> 72

<220>
<223> Zap70

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<400> 72
Met Glu Met Ala Gly Gly Gly Pro Leu His Lys Phe Leu Val Gly Lys
 1          5          10          15
Arg Glu Glu Ile
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$\langle 212 \rangle$	PRT
$\langle 213 \rangle$	unknown

<220>
<223> Jak1

<400> 73
Met Glu Phe Leu Pro Ser Gly Ser Leu Lys Glu Tyr Leu Pro Lys Asn
1 5 10 15

THE

Lys Asn Lys Ile
20

<210> 74
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<212> PRT
<213> unknown

<220>
<223> Jak2

<400> 74

Met	Glu	Tyr	Leu	Pro	Tyr	Gly	Ser	Leu	Arg	Asp	Tyr	Leu	Gln	Lys	His
1				5					10					15	
Lys	Glu	Arg	Ile												
			20												

<210> 75
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<212> PRT
<213> unknown

<220>
<223> Jak3

<400> 75

Met	Glu	Tyr	Leu	Pro	Ser	Gly	Cys	Leu	Arg	Asp	Phe	Leu	Gln	Arg	His
1				5					10					15	
Arg	Ala	Arg	Leu												
			20												

<210> 76
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<220>
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<400> 76

Met	Glu	Tyr	Val	Pro	Leu	Gly	Ser	Leu	Arg	Asp	Tyr	Leu	Pro	Arg	His
1				5					10					15	
Ser	Ile														

<210> 77
<211> 19
<212> PRT
<213> unknown

<220>
<223> Iak1

<400> 77

Leu Glu Tyr Ala Pro Leu Gly Thr Val Tyr Arg Glu Leu Gln Lys Leu
 1 5 10 15
 Ser Lys Phe

<210> 78

<211> 19

<212> PRT

<213> unknown

<220>

<223> Chk1

<400> 78

Leu Glu Tyr Cys Ser Gly Gly Glu Leu Phe Asp Arg Ile Glu Pro Asp
 1 5 10 15
 Ile Gly Met

<210> 79

<211> 22

<212> PRT

<213> unknown

<220>

<223> IKK-1

<400> 79

Met Glu Tyr Cys Ser Gly Gly Asp Leu Arg Lys Leu Leu Asn Lys Pro
 1 5 10 15
 Glu Asn Cys Cys Gly Leu
 20

<210> 80

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<400> 80

Met Glu Tyr Cys Gln Gly Gly Asp Leu Arg Lys Tyr Leu Asn Gln Phe
 1 5 10 15
 Glu Asn Cys Cys Gly Leu
 20

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<212> PRT
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<220>
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<400> 81
 Leu Glu Leu Val Ala Gly Gly Glu Leu Phe Asp Phe Leu Ala Glu Lys
 1 5 10 15
 Glu Ser Leu

<210> 82
 <211> 31
 <212> PRT
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<220>
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<400> 82
 Met Glu Leu Met Ala His Gly Asp Leu Lys Ser Tyr Leu Arg Ser Leu
 1 5 10 15
 Arg Pro Glu Ala Glu Asn Asn Pro Gly Arg Pro Pro Pro Thr Leu
 20 25 30

<210> 83
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 <212> PRT
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<220>
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<400> 83
 Thr Ala Phe His Ala Lys Gly Asn Leu Gln Glu Tyr Leu Thr Arg His
 1 5 10 15
 Val Ile

<210> 84
 <211> 18
 <212> PRT
 <213> unknown

<220>
 <223> ACTRIIA

<400> 84
 Thr Ala Phe His Glu Lys Gly Ser Leu Ser Asp Phe Leu Lys Ala Asn
 1 5 10 15
 Val Val

<210> 85
 <211> 18
 <212> PRT
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<220>
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<400> 85
 Thr Ala Phe His Asp Lys Gly Ser Leu Thr Asp Tyr Leu Lys Gly Asn
 1 5 10 15
 Ile Ile

<210> 86
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 <212> PRT
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<220>
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<400> 86
 Thr His Tyr His Glu His Gly Ser Leu Tyr Asp Phe Leu Gln Arg Gln
 1 5 10 15
 Thr Leu

<210> 87
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 <212> PRT
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<400> 87
 Thr His Tyr His Glu Met Gly Ser Leu Tyr Asp Tyr Leu Gln Leu Thr
 1 5 10 15
 Thr Leu

<210> 88
 <211> 18
 <212> PRT
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<400> 88
 Thr Asp Tyr His Glu Asn Gly Ser Leu Tyr Asp Phe Leu Lys Cys Ala

25/55

1 5 10 15
Thr Leu

<210> 89
<211> 18
<212> PRT
<213> unknown

<220>
<223> ALK4

<400> 89
Ser Asp Tyr His Glu His Gly Ser Leu Phe Asp Tyr Leu Asn Arg Tyr
1 5 10 15
Thr Val

<210> 90
<211> 18
<212> PRT
<213> unknown

<220>
<223> alk6

<400> 90
Thr Asp Tyr His Glu Asn Gly Ser Leu Tyr Asp Tyr Leu Lys Ser Thr
1 5 10 15
Thr Leu

<210> 91
<211> 18
<212> PRT
<213> unknown

<220>
<223> DDR1

<400> 91
Thr Asp Tyr Met Glu Asn Gly Asp Leu Asn Gln Phe Leu Ser Ala His
1 5 10 15
Gln Leu

<210> 92
<211> 18
<212> PRT
<213> unknown

<220>
<223> DDR2

<400> 92

Thr Glu Tyr Met Glu Asn Gly Asp Leu Asn Gln Phe Leu Ser Arg His

1

5

10

15

Glu Pro

<210> 93

<211> 21

<212> PRT

<213> unknown

<220>

<223> ILK

<400> 93

Glu Pro

Thr His Trp Met Pro Tyr Gly Ser Leu Tyr Asn Val Leu His Glu Gly
 1 5 10 15
 Thr Asn Phe Val Val
 20

<210> 94
 <211> 16
 <212> PRT
 <213> unknown

<220>
 <223> JNK

<400> 94
 Met Glu Leu Met Asp Ala Asn Leu Cys Gln Val Ile Gln Met Glu Leu
 1 5 10 15

<210> 95
 <211> 20
 <212> PRT
 <213> Artificial Sequence

<220>
 <221> MYRISTATE
 <222> (1)...(0)
 <223>

<221> AMIDATION
 <222> (0)...(20)

<223> Akt1/Raca

<400> 95
 Gly Met Glu Tyr Ala Asn Gly Gly Glu Leu Phe Phe His Leu Ser Arg
 1 5 10 15
 Glu Arg Val Phe
 20

<210> 96
 <211> 19
 <212> PRT
 <213> Artificial Sequence

<220>
 <221> MYRISTATE
 <222> (1)...(0)

<221> AMIDATION
 <222> (0)...(19)

<223> Alk1

<400> 96
 Gly Thr His Tyr His Glu His Gly Ser Leu Tyr Asp Phe Leu Gln Arg
 1 5 10 15

Gln Thr Leu

<210> 97
 <211> 22
 <212> PRT
 <213> Artificial Sequence

<220>
 <221> ACETYLATION
 <222> (1)...(0)

<221> AMIDATION
 <222> (0)...(22)

<223> Braf

<400> 97
 Lys Lys Lys Lys Lys Lys Gly Gly Ser Ser Leu Tyr His His Leu His
 1 5 10 15
 Ile Ile Glu Thr Lys Phe
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<210> 98
 <211> 21
 <212> PRT
 <213> Artificial Sequence

<220>
 <221> MYRISTATE
 <222> (1)...(0)

<221> AMIDATION
 <222> (0)...(21)

<223> Braf

<400> 98
 Gly Thr Gln Trp Ser Glu Gly Ser Ser Leu Tyr His His Leu His Ile
 1 5 10 15
 Ile Glu Thr Lys Phe
 20

<210> 99
 <211> 22
 <212> PRT
 <213> Artificial Sequence

<220>
 <221> MYRISTATE
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<221> AMIDATION
 <222> (0)...(22)

<223> c-Ab1

Cys Asn Arg Gln Glu Val
20

<213> Artificial Sequence

 $\langle 222 \rangle \quad (1) \dots (0)$

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<221> AMIDATION
<222> (0) ... (21)
<223>

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Glu Thr His Asn Pro
20

Time (s)	Angle (°)	Intensity (a.u.)	Index	Phase (°)	Amplitude (a.u.)	Frequency (Hz)	Wavelength (nm)	Order
0.0	0.0	1.0	<210>	101	1.0	1.0	1.0	1
0.1	0.1	0.5	<211>	21	0.5	0.5	0.5	1
0.2	0.2	0.1	<212>	PRT	0.1	0.1	0.1	1
0.3	0.3	0.0	<213>	Artificial Sequence	0.0	0.0	0.0	1

$\langle 222 \rangle \quad (1) \dots (0)$

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<221> AMIDATION
<222> (0) ... (21)
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Gln Glu Thr Lys Phe
20

<213> Artificial Sequence

$\langle 222 \rangle \quad (1) \dots (0)$

<223> benzyl ester at position 11

<221> AMIDATION

<222> (0)...(14)

<223> c-Raf

<400> 102

Ser Ser Leu Tyr Lys His Leu His Val Gln Glu Thr Lys Phe
1 5 10

<210> 103

<211> 21

<212> PRT

<213> Artificial Sequence

<220>

<221> MYRISTATE

<222> (1)...(0)

<221> AMIDATION

<222> (0)...(21)

<223> c-Sea

<400> 103

Gly Leu Pro Tyr Met Arg His Gly Asp Leu Arg His Phe Ile Arg Ala
1 5 10 15
Gln Glu Arg Ser Pro
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<210> 104

<211> 22

<212> PRT

<213> Artificial Sequence

<220>

<221> MYRISTATE

<222> (1)...(0)

<221> AMIDATION

<222> (0)...(22)

<223> c-Src

<400> 104

Gly Thr Glu Tyr Met Ser Lys Gly Ser Leu Leu Asp Phe Leu Lys Gly
1 5 10 15
Glu Thr Gly Lys Tyr Leu
20

<210> 105

<211> 14

<212> PRT

<213> Artificial Sequence

<220>
 <221> ACETYLATION
 <222> (1)...(0)
 <223> benzyl ester at position 5
 benzyl ester at position 9

 <221> AMIDATION
 <222> (0)...(14)

 <223> c-Src

 <400> 105
 Gly Ser Leu Leu Asp Leu Lys Gly Glu Thr Gly Lys Phe Leu
 1 5 10

<210> 106
 <211> 21
 <212> PRT
 <213> Artificial Sequence

<220>
 <221> MYRISTATE
 <222> (1)...(0)

 <221> AMIDATION
 <222> (0)...(21)
 <223>

<223> CDK2

<400> 106
 Gly Phe Glu Phe Leu His Gln Asp Leu Lys Lys Phe Met Asp Ala Ser
 1 5 10 15
 Ala Leu Thr Gly Ile
 20

<210> 107
 <211> 14
 <212> PRT
 <213> Artificial Sequence

<220>
 <221> ACETYLATION
 <222> (1)...(0)
 <223> benzyl ester at position 1
 benzyl ester at position 7

<221> AMIDATION
 <222> (0)...(14)
 <223>

<223> CDK2

<400> 107

Asp Leu Lys Lys Phe Met Asp Ala Ser Ala Leu Thr Gly Met
 1 5 10

<210> 108
 <211> 14
 <212> PRT
 <213> Artificial Sequence

<220>
 <221> ACETYLATION
 <222> (1)...(0)
 <223> benzyl ester at position 1
 benzyl ester at position 7

<221> AMIDATION
 <222> (0)...(14)

<223> CDK4

<400> 108

Asp Leu Arg Thr Tyr Leu Asp Lys Ala Pro Pro Pro Gly Leu
 1 5 10

<210> 109
 <211> 21
 <212> PRT
 <213> Artificial Sequence

<220>
 <221> MYRISTATE
 <222> (1)...(0)

<221> AMIDATION
 <222> (0)...(21)

<223> CDK4

<400> 109

Gly Phe Glu His Val Asp Gln Asp Leu Arg Thr Tyr Leu Asp Lys Ala
 1 5 10 15
 Pro Pro Pro Gly Leu
 20

<210> 110
 <211> 21
 <212> PRT
 <213> Artificial Sequence

<220>
 <221> MYRISTATE
 <222> (1)...(0)

<221> AMIDATION
 <222> (0)...(21)

<223> CDK6

<400> 110

Gly	Phe	Glu	His	Val	Asp	Gln	Asp	Leu	Thr	Thr	Tyr	Leu	Asp	Lys	Val
1				5					10					15	
Pro	Glu	Pro	Gly	Val											
				20											

<210> 111

<211> 19

<212> PRT

<213> Artificial Sequence

<220>

<221> MYRISTATE

<222> (1)...(0)

<221> AMIDATION

<222> (0)...(19)

<223> Chk1

<400> 111

Gly	Glu	Tyr	Ser	Ser	Gly	Gly	Glu	Leu	Phe	Asp	Arg	Ile	Glu	Pro	Asp
1				5					10					15	
Ile	Gly	Met													

<210> 112

<211> 19

<212> PRT

<213> Artificial Sequence

<220>

<221> MYRISTATE

<222> (1)...(0)

<221> AMIDATION

<222> (0)...(19)

<223>

<223> Chk1

<400> 112

Gly	Glu	Tyr	Ala	Ser	Gly	Gly	Glu	Leu	Phe	Asp	Arg	Ile	Glu	Pro	Asp
1				5					10					15	
Ile	Gly	Met													

<210> 113

<211> 19

<212> PRT

<213> Artificial Sequence

<220>

<221> ACETYLATION

<222> (1)...(0)

<221> AMIDATION
 <222> (0)...(19)
 <223>

<223> CK IIa

<400> 113

Lys	Lys	Lys	Lys	Lys	Gly	Gly	Asn	Asn	Thr	Asp	Phe	Lys	Gln	Leu	Tyr
1				5					10					15	
Gln	Thr	Leu													

<210> 114

<211> 17

<212> PRT

<213> Artificial Sequence

<220>

<221> MYRISTATE

<222> (1)...(0)

<221> AMIDATION

<222> (0)...(17)

<223> CK IIa

<400> 114

Gly	Phe	Glu	His	Val	Asn	Asn	Thr	Asp	Phe	Lys	Gln	Leu	Tyr	Gln	Thr
1				5					10					15	
Leu															

<210> 115

<211> 22

<212> PRT

<213> Artificial Sequence

<220>

<221> MYRISTATE

<222> (1)...(0)

<221> AMIDATION

<222> (0)...(22)

<223>

<223> Csk

<400> 115

Gly	Thr	Glu	Tyr	Met	Ala	Lys	Gly	Ser	Leu	Val	Asp	Tyr	Leu	Arg	Ser
1				5					10					15	
Arg	Gly	Arg	Ser	Val	Leu										
				20											

<210> 116

<211> 14
 <212> PRT
 <213> Artificial Sequence

<220>
 <221> ACETYLTATION
 <222> (1)...(0)
 <223> benzyl ester at position 5

<221> AMIDATION
 <222> (0)...(14)

<223> Csk

<400> 116
 Gly Ser Leu Val Asp Leu Arg Ser Arg Gly Arg Ser Val Leu
 1 5 10

<210> 117
 <211> 21
 <212> PRT
 <213> Artificial Sequence

<220>
 <221> MYRISTATE
 <222> (1)...(0)

<221> AMIDATION
 <222> (0)...(21)

<223> Fak

<400> 117
 Gly Met Glu Leu Ser Thr Leu Gly Glu Leu Arg Ser Phe Leu Gln Val
 1 5 10 15
 Arg Lys Tyr Ser Leu
 20

<210> 118
 <211> 17
 <212> PRT
 <213> Artificial Sequence

<220>
 <221> MYRISTATE
 <222> (1)...(0)

<221> AMIDATION
 <222> (0)...(17)

<223> FGFR-3

<400> 118
 Gly Gly Asn Leu Arg Glu Phe Leu Arg Ala Arg Arg Pro Pro Gly Leu
 1 5 10 15
 Glu

<210> 119
 <211> 16
 <212> PRT
 <213> Artificial Sequence

<220>
 <221> ACETYLATION
 <222> (1)...(0)
 <223> benzyl ester at position 5
 benzyl ester at position 16

<221> AMIDATION
 <222> (0)...(16)

<223> FGFR-3

COOT
 <400> 119
 Gly Asn Leu Arg Glu Phe Leu Arg Ala Arg Arg Pro Pro Gly Leu Glu
 1 5 10 15

<210> 120
 <211> 23
 <212> PRT
 <213> Artificial Sequence

<220>
 <221> MYRISTATE
 <222> (1)...(0)
 <221> AMIDATION
 <222> (0)...(23)

<223> FGFR-3

<400> 120
 Gly Val Glu Tyr Ala Ala Lys Gly Asn Leu Arg Glu Phe Leu Arg Ala
 1 5 10 15
 Arg Arg Pro Pro Gly Leu Glu
 20

<210> 121
 <211> 13
 <212> PRT
 <213> Artificial Sequence

<220>
 <223> stearyl at position 1

<221> AMIDATION
 <222> (0)...(13)
 <223> FGFR-3

<400> 121
 Gly Ser Phe Asp Thr Ser Lys Pro Pro Glu Glu Gln Leu
 1 5 10

<210> 122
 <211> 23
 <212> PRT
 <213> Artificial Sequence

<220>
 <221> MYRISTATE
 <222> (1)...(0)

<221> AMIDATION
 <222> (0)...(23)

<223> Flk1

<400> 122
 Gly Val Glu Phe Ser Lys Phe Gly Asn Leu Ser Asn Phe Leu Arg Ala
 1 5 10 15
 Lys Arg Asn Leu Phe Val Pro
 20

<210> 123
 <211> 17
 <212> PRT
 <213> Artificial Sequence

<220>
 <221> MYRISTATE
 <222> (1)...(0)
 <221> AMIDATION
 <222> (0)...(17)
 <223>

<223> Flk1

<400> 123
 Gly Gly Asn Leu Ser Asn Phe Leu Arg Ala Lys Arg Asn Leu Phe Val
 1 5 10 15
 Pro

<210> 124
 <211> 16
 <212> PRT
 <213> Artificial Sequence

<220>
 <221> ACETYLATION
 <222> (1)...(0)

<221> AMIDATION
 <222> (0)...(16)

<223> Flk1

<400> 124

Gly	Asn	Leu	Ser	Asn	Phe	Leu	Arg	Ala	Lys	Arg	Asn	Leu	Phe	Val	Pro
1				5					10				15		

<210> 125

<211> 13

<212> PRT

<213> Artificial Sequence

<220>

<223> stearyl at position 1

<221> AMIDATION

<222> (0)...(13)

<223> Flk1

<400> 125

Gly	Arg	Phe	Arg	Gln	Gly	Lys	Asp	Tyr	Val	Gly	Glu	Leu
1				5					10			

<210> 126

<211> 22

<212> PRT

<213> Artificial Sequence

<220>

<221> ACETYLATION

<222> (1)...(0)

<221> AMIDATION

<222> (0)...(22)

<223> GSK3b

<400> 126

Lys	Lys	Lys	Lys	Lys	Lys	Lys	Gly	Gly	Gly	Val	Ala	Arg	His	Tyr	Ser	Arg
1				5						10					15	
Ala	Lys	Gln	Thr	Leu	Pro											
			20													

<210> 127

<211> 13

<212> PRT

<213> Artificial Sequence

<220>

<221> ACETYLATION

<222> (1)...(0)

<221> AMIDATION

<222> (0)...(13)

<223> GSK3b

<400> 127

Val Ala Arg His Tyr Ser Arg Ala Lys Gln Thr Leu Pro
1 5 10

<210> 128

<211> 22

<212> PRT

<213> Artificial Sequence

<220>

<221> MYRISTATE

<222> (1)...(0)

<221> AMIDATION

<222> (0)...(22)

<223> GSK3b

<400> 128

E08070-CT38E07

40/55

Gly Asp Tyr Val Pro Glu Thr Val Tyr Arg Val Ala Arg His Tyr Ser
1 5 10 15
Arg Ala Lys Gln Thr Leu
20

<210> 129
<211> 12
<212> PRT
<213> Artificial Sequence

<220>
<221> ACETYLATION
<222> (1)...(0)

<221> AMIDATION
<222> (0)...(12)

<223> GSK3b

<400> 129
Arg Val Ala Arg His Tyr Ser Arg Ala Lys Gln Thr
1 5 10

<210> 130
<211> 22
<212> PRT
<213> Artificial Sequence

<220>
<221> MYRISTATE
<222> (1)...(0)

<221> AMIDATION
<222> (0)...(22)

<223> Hck

<400> 130
Gly Thr Glu Phe Met Ala Lys Gly Ser Leu Leu Asp Phe Leu Lys Ser
1 5 10 15
Asp Glu Gly Ser Lys Gln
20

<210> 131
<211> 20
<212> PRT
<213> Artificial Sequence

<220>
<221> MYRISTATE
<222> (1)...(0)

<221> AMIDATION

<222> (0)...(20)

<223> Iak1

<400> 131

Gly Leu Glu Tyr Ala Pro Leu Gly Thr Val Tyr Arg Glu Leu Gln Lys
 1 5 10 15
 Leu Ser Lys Phe
 20

<210> 132

<211> 23

<212> PRT

<213> Artificial Sequence

<220>

<221> MYRISTATE

<222> (1)...(0)

<221> AMIDATION

<222> (0)...(23)

<223> IKK-1

<400> 132

Gly Met Glu Tyr Ser Ser Gly Gly Asp Leu Arg Lys Leu Leu Asn Lys
 1 5 10 15
 Pro Glu Asn Ser Ser Gly Leu
 20

<210> 133

<211> 23

<212> PRT

<213> Artificial Sequence

<220>

<221> MYRISTATE

<222> (1)...(0)

<221> AMIDATION

<222> (0)...(23)

<223>

<223> IKK-2

<400> 133

Gly Met Glu Tyr Ser Gln Gly Gly Asp Leu Arg Lys Tyr Leu Asn Gln
 1 5 10 15
 Phe Glu Asn Ser Ser Gly Leu
 20

<210> 134
 <211> 22
 <212> PRT
 <213> Artificial Sequence

<220>
 <221> MYRISTATE
 <222> (1)...(0)

<221> AMIDATION
 <222> (0)...(22)

<223> ILK

<400> 134
 Gly Thr His Trp Met Pro Tyr Gly Ser Leu Tyr Asn Val Leu His Glu
 1 5 10 15
 Gly Thr Asn Phe Val Val
 20

<210> 135
 <211> 13
 <212> PRT
 <213> Artificial Sequence

<220>
 <223> stearyl at position 1

<221> AMIDATION
 <222> (0)...(13)
 <223> ILK

<400> 135
 Gly Tyr Asn Val Leu His Glu Gly Thr Asn Phe Val Val
 1 5 10

<210> 136
 <211> 19
 <212> PRT
 <213> Artificial Sequence

<220>
 <221> MYRISTATE
 <222> (1)...(0)

<221> AMIDATION
 <222> (0)...(19)
 <223>

<223> IRK

<400> 136

Gly Met Glu Leu Met Ala His Gly Asp Leu Lys Ser Tyr Leu Arg Ser
 1 5 10 15
 Leu Arg Pro

<210> 137

<211> 12

<212> PRT

<213> Artificial Sequence

<220>

<221> ACETYLATION

<222> (1)...(0)

<221> AMIDATION

<222> (0)...(12)

<223> IRK

<400> 137

Ala Gln Asn Asn Pro Gly Arg Pro Pro Pro Thr Leu
 1 5 10

<210> 138

<211> 13

<212> PRT

<213> Artificial Sequence

<220>

<221> MYRISTATE

<222> (1)...(0)

<221> AMIDATION

<222> (0)...(13)

<223> IRK

<400> 138

Gly Leu Lys Ser Tyr Leu Arg Ser Leu Arg Pro Glu Ala
 1 5 10

<210> 139

<211> 13

<212> PRT

<213> Artificial Sequence

<220>

<221> MYRISTATE

<222> (1)...(0)

<221> AMIDATION

<222> (0)...(13)

<223> IRK

<400> 139

Gly Ala Glu Asn Asn Pro Gly Arg Pro Pro Pro Thr Leu
 1 5 10

<210> 140

<211> 17

<212> PRT

<213> Artificial Sequence

<220>

<221> MYRISTATE

<222> (1)...(0)

<221> AMIDATION

<222> (0)...(17)

<223> IRK

<400> 140

Gly Leu Arg Pro Glu Ala Glu Asn Asn Pro Gly Arg Pro Pro Pro Thr
 1 5 10 15
 Leu

<210> 141

<211> 21

<212> PRT

<213> Artificial Sequence

<220>

<221> MYRISTATE

<222> (1)...(0)

<221> AMIDATION

<222> (0)...(21)

<223> Jak1

<400> 141

Gly Met Glu Phe Leu Pro Ser Gly Ser Leu Lys Glu Tyr Leu Pro Lys
 1 5 10 15
 Asn Lys Asn Lys Ile
 20

<210> 142

<211> 13

<212> PRT

<213> Artificial Sequence

<220>

<221> MYRISTATE

<222> (1)...(0)

<221> AMIDATION

<222> (0)...(13)

<223> Jak1

<400> 142

Gly Leu Lys Glu Tyr Leu Pro Lys Asn Lys Asn Lys Ile
1 5 10

<210> 143

<211> 13

<212> PRT

<213> Artificial Sequence

<220>

<221> MYRISTATE

<222> (1)...(0)

<221> AMIDATION

<222> (0)...(13)

<223> Jak2

<400> 143

Gly Leu Arg Asp Tyr Leu Gln Lys His Lys Glu Arg Ile
1 5 10

<210> 144

<211> 11

<212> PRT

<213> Artificial Sequence

<220>

<223> stearyl at position 1

<221> AMIDATION

<222> (0)...(11)

<223> Jak2

<400> 144

Gly Leu Arg Asp Tyr Leu Gln Lys His Lys Glu
1 5 10

<210> 145

<211> 20

<212> PRT

<213> Artificial Sequence

<220>

<221> MYRISTATE

<222> (1)...(0)

<221> AMIDATION

<222> (0)...(20)

1. The first step is to identify the problem. This involves understanding the symptoms and the context in which they are occurring.

Gly Met Glu Tyr Leu Pro Ser Gly Ser Leu Arg Asp Phe Leu Gln Arg
1 5 10 15
His Arg Ala Leu
20

<213> Artificial Sequence

<222> (0) . . . (21)

Gly Met Glu Tyr Leu Pro Ser Gly Ser Leu Arg Asp Phe Leu Gln Arg
1 5 10 15
His Arg Ala Arg Leu
20

<213> Artificial Sequence

<222> (0) ... (13)

Gly Leu Arg Asp Phe Leu Gln Arg His Arg Ala Arg Leu
1 5 10

<213> Artificial Sequence

<221> ACETYLATION

<222> (1)...(0)
 <223> benzyl ester at position 5

<221> AMIDATION
 <222> (0)...(14)

<223> Lck

<400> 148

Gly Ser Leu Val Asp Leu Lys Thr Pro Ser Gly Ile Lys Leu
 1 5 10

<210> 149

<211> 22

<212> PRT

<213> Artificial Sequence

<220>

<221> MYRISTATE

<222> (1)...(0)

<221> AMIDATION

<222> (0)...(22)

<223> Lck

<400> 149

Gly Thr Glu Tyr Met Glu Asn Gly Ser Leu Val Asp Phe Leu Lys Thr
 1 5 10 15
 Pro Ser Gly Ile Lys Leu
 20

<210> 150

<211> 22

<212> PRT

<213> Artificial Sequence

<220>

<221> MYRISTATE

<222> (1)...(0)

<221> AMIDATION

<222> (0)...(22)

<223> Lyn

<400> 150

Gly Thr Glu Tyr Met Ala Lys Gly Ser Leu Leu Asp Phe Leu Lys Ser
 1 5 10 15
 Asp Glu Gly Gly Lys Val
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<210> 151

<211> 20

<212> PRT

<213> Artificial Sequence

<220>

<221> MYRISTATE

<222> (1)...(0)

<221> AMIDATION

<222> (0)...(20)

<223> MARK1

<400> 151

Gly Met Glu Tyr Ala Ser Gly Gly Glu Val Phe Asp Tyr Leu Val Ala

1 5 10 15

His Gly Arg Met

20

<210> 152

<211> 15

<212> PRT

<213> Artificial Sequence

<220>

<221> ACETYLATION

<222> (1)...(0)

<223> benzyl ester at position 2

benzyl ester at position 5

<221> AMIDATION

<222> (0)...(15)

<223> PDGFR-b

<400> 152

Gly Asp Leu Val Asp Tyr Leu His Arg Asn Lys His Thr Phe Leu

1 5 10 15

<210> 153

<211> 22

<212> PRT

<213> Artificial Sequence

<220>

<221> MYRISTATE

<222> (1)...(0)

<221> AMIDATION

<222> (0)...(22)

<223> PDGFR-b

<400> 153

Gly Thr Glu Tyr Ser Arg Tyr Gly Asp Leu Val Asp Tyr Leu His Arg

1 5 10 15

Asn Lys His Thr Phe Leu

20

<210> 154
 <211> 20
 <212> PRT
 <213> Artificial Sequence

<220>
 <221> MYRISTATE
 <222> (1)...(0)

<221> AMIDATION
 <222> (0)...(20)

<223> PKCb

<400> 154

Gly	Met	Glu	Tyr	Val	Asn	Gly	Gly	Asp	Leu	Met	Tyr	His	Ile	Gln	Gln
1				5					10					15	
Val	Gly	Arg	Phe												
			20												

<210> 155
 <211> 20
 <212> PRT
 <213> Artificial Sequence

<220>
 <221> ACETYLTATION
 <222> (1)...(0)

<221> AMIDATION
 <222> (0)...(20)

<223> PKCb

<400> 155

Lys	Lys	Lys	Lys	Lys	Lys	Gly	Gly	Asp	Leu	Met	Tyr	His	Ile	Gln	Gln
1				5					10					15	
Val	Gly	Arg	Phe												
			20												

<210> 156
 <211> 12
 <212> PRT
 <213> Artificial Sequence

<220>
 <221> ACETYLTATION
 <222> (1)...(0)
 <223> benzyl ester at position 5

<221> AMIDATION
 <222> (0)...(12)

<223> Plk

<400> 156

Arg Ser Leu Leu Glu Leu His Lys Arg Arg Lys Ala

1

5

10

<210> 157
 <211> 13
 <212> PRT
 <213> Artificial Sequence

<220>
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<221> AMIDATION
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<223> Plk

<400> 157
 Gly Arg Ser Leu Leu Glu Leu His Lys Arg Arg Lys Ala
 1 5 10

<210> 158
 <211> 20
 <212> PRT
 <213> Artificial Sequence

<220>
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 <221> AMIDATION
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<223> Plk

<400> 158
 Gly Leu Glu Leu Ser Arg Arg Arg Ser Leu Leu Glu Leu His Lys Arg
 1 5 10 15
 Arg Lys Ala Leu
 20

<210> 159
 <211> 22
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<220>
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<221> AMIDATION
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<223> Ret

<400> 159

Gly Val Glu Tyr Ala Lys Tyr Gly Ser Leu Arg Gly Phe Leu Arg Glu
 1 5 10 15
 Ser Arg Lys Val Gly Pro
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<210> 160
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 <223> benzyl ester at position 9

<221> AMIDATION
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<223> Ret

<400> 160

Gly Ser Leu Arg Gly Phe Leu Arg Glu Ser Arg Lys Val Gly Pro
 1 5 10 15

<210> 161
 <211> 21
 <212> PRT
 <213> Artificial Sequence

<220>
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<221> AMIDATION
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<223> Ron

<400> 161

Gly Leu Pro Tyr Met Cys His Gly Asp Leu Leu Gln Phe Ile Arg Ser
 1 5 10 15
 Pro Gln Arg Asn Pro
 20

<210> 162
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<220>
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 <222> (1)...(0)

<221> AMIDATION
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<223> SNK

<400> 162

Gly Leu Glu Tyr Ser Ser Arg Arg Ser Met Ala His Ile Leu Lys Ala
 1 5 10 15
 Arg Lys Val Leu
 20

<210> 163

<211> 20

<212> PRT

<213> Artificial Sequence

<220>

<221> MYRISTATE

<222> (1)...(0)

<221> AMIDATION

<222> (0)...(20)

<223> Syk

<400> 163

Gly Met Glu Met Ala Glu Leu Gly Pro Leu Asn Lys Tyr Leu Gln Gln
 1 5 10 15
 Asn Arg His Val
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<210> 164

<211> 19

<212> PRT

<213> Artificial Sequence

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<221> MYRISTATE

<222> (1)...(0)

<221> AMIDATION

<222> (0)...(19)

<223> TGFbRII

<400> 164

Gly Thr Ala Phe His Ala Lys Gly Asn Leu Gln Glu Tyr Leu Thr Arg
 1 5 10 15
 His Val Ile

<210> 165

<211> 25

<212> PRT

<213> Artificial Sequence

<220>

<221> MYRISTATE

<222> (1)...(0)

<221> AMIDATION
 <222> (0)...(25)

<223> TrkB

<400> 165
 Gly Phe Glu Tyr Met Lys His Gly Asp Leu Asn Lys Phe Leu Arg Ala
 1 5 10 15
 His Gly Pro Asp Ala Val Leu Met Ala
 20 25

<210> 166
 <211> 13
 <212> PRT
 <213> Artificial Sequence

<220>
 <221> MYRISTATE
 <222> (1)...(0)
 <221> AMIDATION
 <222> (0)...(13)

<223> TrkB

<400> 166
 Gly Leu Arg Ala His Gly Pro Asp Ala Val Leu Met Ala
 1 5 10

<210> 167
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<220>
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 <222> (1)...(0)
 <221> AMIDATION
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<223> TrkB

<400> 167
 Gly Leu Arg Ala His Gly Pro Asp Ala Val Leu
 1 5 10

<210> 168
 <211> 13
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 <221> MYRISTATE
 <222> (1)...(0)

<221> AMIDATION
 <222> (0)...(13)

<223> TrkB

<400> 168
 Gly Leu Asn Phe Lys Leu Arg Ala His Gly Pro Asp Ala
 1 5 10

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 <211> 13
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 <222> (0)...(13)

<223> TrkB

<400> 169
 Gly Phe Lys Leu Arg Ala His Gly Pro Asp Ala Val Leu
 1 5 10

<210> 170
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 <222> (1)...(0)

<221> AMIDATION
 <222> (0)...(21)

<223> Zap70

<400> 170
 Gly Met Glu Met Ala Gly Gly Gly Pro Leu His Lys Phe Leu Val Gly
 1 5 10 15
 Lys Arg Glu Glu Ile
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<210> 171
 <211> 21
 <212> PRT
 <213> Unknown

<220>

<223> IRK

<400> 171

Met Ala His Gly Asp Leu Lys Ser Tyr Leu Arg Ser Leu Arg Pro Glu
 1 5 10 15
 Ala Glu Asn Asn Pro
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<210> 172

<211> 8

<212> PRT

<213> Unknown

<220>

<223> endothelial growth factor receptor

<400> 172

Lys Phe Asp Val Ile Asn Leu Ala
 1 5

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